In the Claims:

1-118. Canceled.

- 119. (Currently amended) An isolated polypeptide having at least 80% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207);
- (b) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135 (SEQ ID NO: 207);
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135(SEQ ID NO: 207), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951,

 wherein the nucleic acid encoding said polypeptide is amplified in lung or colon tumor.
- 120. (Currently amended) The isolated polypeptide of Claim 39 having at least 85% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207);
- (b) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 SEQ ID NO: 207), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135 (SEQ ID NO: 207);
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135(SEQ ID NO: 207), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951, wherein the nucleic acid encoding said polypeptide is amplified in lung or colon tumor.

- 121. (Currently amended) The isolated polypeptide of Claim 39 having at least 90% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207);
- (b) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135 (SEO ID NO: 207);
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135(SEQ ID NO: 207), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951,
 wherein the nucleic acid encoding said polypeptide is amplified in lung or colon tumor.
- 122. (Currently amended) The isolated polypeptide of Claim 39 having at least 95% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207);
- (b) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135 (SEQ ID NO: 207);
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135(SEQ ID NO: 207), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951,
 wherein the nucleic acid encoding said polypeptide is amplified in lung or colon tumor.
- 123. (Currently amended) The isolated polypeptide of Claim 39 having at least 99% amino acid sequence identity to:

- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207);
- (b) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207), lacking its associated signal peptide,
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135 (SEQ ID NO: 207);
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135(SEQ ID NO: 207), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951, wherein the nucleic acid encoding said polypeptide is amplified in lung or colon tumor.
- 124. (Currently amended) An isolated polypeptide comprising:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207);
- (b) the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135 (SEQ ID NO: 207);
- (d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 135(SEQ ID NO: 207), lacking its associated signal peptide; or
- the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951;
 wherein the nucleic acid encoding said polypeptide is amplified in lung or colon tumor.
- 125. (Currently amended) The isolated polypeptide of Claim 124 comprising the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207).
- 126. (Currently amended) The isolated polypeptide of Claim 124 comprising the amino acid sequence of the polypeptide of SEQ ID NO: 207 shown in Figure 135 (SEQ ID NO: 207), lacking its associated signal peptide.

127-128. Canceled.

- 129. (Previously presented) The isolated polypeptide of Claim 124 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209951.
- 130. (Currently amended) A chimeric polypeptide comprising a polypeptide according to Claim 124 119 fused to a heterologous polypeptide.
- 131. (Previously presented) The chimeric polypeptide of Claim 130, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.